

SEQUENCE LISTING

<110> National Agricultural Research Organization; Katsuji Watanabe

<120> METHOD AND SYSTEM FOR SEARCHING FOR RELATIONSHIPS
BETWEEN BASE SEQUENCE IN GENES

<160>4

<210>1

<211>1009

<212>DNA

<213>Rhodospirillum rubrum

<220>

<221>rRNA

<222>(50),(51),(63),(463),(468),(574),(687),(806),(853),(854),(888)

,(889),(966)

<223>unknown

<400>

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121 ggaaacggca gctaataccg gatacgccgt attgggaaag aaattcggcc ttggatgggc

181 ccgcgttggg ttagctagat ggtggggtaa cgccctacca tggcgacgat ccatagctgg

241 ttgagagga tgatcagcca cactgggact gagacacggc ccagactcct acgggaggca

301 gcagtgggga atcttagaca atgggggcaa ccctgatcta gccatgccg gtgagtgatg

361 aaggccttag ggttgtaaag ctcttcagc aggggaagata atgactgtac ctgcagaaga

421 agctccggct aactccgtgc cagcagccgc ggtaatacgg agnnggcnag cgtgttcggg

481 aattactggg cgtaaagcgc gcgtaggcgg atcggtcagt tgggggtgaa agccccgggc

541 tcaacctcgg aactgccctc aaaactaccg atcnagagtt cgggagaggt aagcggaatt

601 cccagtgtag aggtgaaatt cgtagatatt gggagaaca ccagtggcga aggcggctta
 661 ctggaccgat actgacgctg aggtgcnaaa gcgtggggag caaacaggat tagataccct
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 841 atgacgggg gcnngcacaa gcggtggagc atgtggttta attcgaanna acgcgcagaa
 901 ccttaccagc tcttgacatc ccgggacgac ttccagagat ggaatttttc acttcgggtga
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<212>DNA

<213>Rhodospirillum salexigens

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<221>rRNA

<222>(1),(72),(73),(85),(485),(490),(596),(709),(828),(875),(876),(
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<223>unknown

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 121 icaggctctt gggataactg ctggaaacgg cagetaatac cggatagccc gtattgggaa
 181 agaaattcgg ccttgatgg gcccgcttg gattagctag atgggtgggt aacgcctac
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421 taatgactgt acctgcagaa gaagctccgg ctaactccgt gccacgagcc gcggtataac
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 1321 acgtcccggt gnnnltgaca caccgccgt cacacatgg gagttggtt gacccgaaga
 1381 cgttgagcta acccgaaagg ggggcagncg gccacggtca ggtcagcgac tggggtnnnn
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<210>3

<211>19

<212>DNA

<213>Artificial Sequence

<220>

<223>Artificial Sequence

<400>

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<210>4

<211>16

<212>DNA

<213>Artificial Sequence

<220>

<223>Artificial Sequence

<400>

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09900875.071001